

<!--StartFragment-->RESULT 2 S7  
AX280095  
LOCUS AX280095 20 bp DNA linear PAT 02-NOV-2001  
DEFINITION Sequence 24 from Patent WO0177295.  
ACCESSION AX280095  
VERSION AX280095.1 GI:16607537  
KEYWORDS .  
SOURCE Escherichia coli  
ORGANISM Escherichia coli  
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;  
Enterobacteriaceae; Escherichia.  
REFERENCE 1  
AUTHORS Denning,D.W., Brookman,J.L., Rickers,A. and Birch,M.  
TITLE Mutant bank  
JOURNAL Patent: WO 0177295-A 24 18-OCT-2001;  
F2G Limited (GB)  
FEATURES Location/Qualifiers  
source 1. .20  
/organism="Escherichia coli"  
/mol\_type="unassigned DNA"  
/db\_xref="taxon:562"  
ORIGIN  
  
Query Match 67.3%; . Score 14.8; DB 6; Length 20;  
Best Local Similarity 88.9%; Pred. No. 1.8e+04;  
Matches 16; Conservative 0; Mismatches 2; Indels 0; Gaps 0;  
  
Qy 5 GACTTGCTGTTTTTGCTC 22  
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Db 1 GCCTTCCTGTTTTTGCTC 18  
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<!--StartFragment-->RESULT 14

AR258547

LOCUS AR258547 40 bp DNA linear PAT 20-DEC-2002

DEFINITION Sequence 36 from patent US 6489146.

ACCESSION AR258547

VERSION AR258547.1 GI:27308923

KEYWORDS .

SOURCE Unknown.

ORGANISM Unknown.

Unclassified.

REFERENCE 1 (bases 1 to 40)

AUTHORS Stemmer;W.P.C. and Lipshutz,R.J.

TITLE End-complementary polymerase reaction

JOURNAL Patent: US 6489146-A 36 03-DEC-2002;

FEATURES Location/Qualifiers

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/organism="unknown"

/mol\_type="genomic DNA"

ORIGIN

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Best Local Similarity 88.9%; Pred. No. 1.9e+04;

Matches 16; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

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Db 7 GCCTTCCTGTTTTTGCTC 24

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12

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<!--StartFragment-->RESULT 9
US-10-257-213-24
; Sequence 24, Application US/10257213
; Publication No. US20030129733A1
; GENERAL INFORMATION:
; APPLICANT: DENNING, David Wemys
; APPLICANT: BROOKMAN, Jayne Louise
; APPLICANT: RICKERS, Andre
; APPLICANT: BIRCH, Mike
; TITLE OF INVENTION: MUTANT BANK
; FILE REFERENCE: 31229-183869
; CURRENT APPLICATION NUMBER: US/10/257,213
; CURRENT FILING DATE: 2002-10-10
; PRIOR APPLICATION NUMBER: GB0008748.6 PCT/GB01/01626
; PRIOR FILING DATE: 2000-04-11
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 24
; LENGTH: 20
; TYPE: DNA
; ORGANISM: Escherichia coli
US-10-257-213-24

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Query Match          67.3%; Score 14.8; DB 15; Length 20;
Best Local Similarity 88.9%; Pred. No. 4.6e+03;
Matches 16; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

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Db          1 GCCTTCCTGTTTTTGCTC 18

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